

Amendments to the Specification:

Please amend the specification as follows:

Page 10, last paragraph (lines 36-37), continuing on page 11 (lines 1-8)

~~Figure 4 shows~~ Figures 4A-C show the comparison of deduced amino acid sequence of wSBE I-D4 cDNA with the deduced amino acid sequence of rice SBE I (RSBE I; Nakamura *et al*, 1992), maize SBE I (MSBE I; Baba *et al*, 1991), wSBE I-D2 type cDNA (D2 cDNA; Rahman *et al*, 1997), pea SBE II (PESBE II, homologous to maize SBE I; Burton *et al*, 1995), and potato SBE I (POSBE; Cangiano *et al*, 1993). The deduced amino acid sequence of the wSBE I-D4 cDNA is denoted by "D4cDNA". Residues present in at least three of the sequences are identified in the consensus sequence in capitals.

Page 11, fourth full paragraph (lines 26-27)

~~Figure 7 shows~~ Figures 7A-B show the hybridisation of SBE I genomic clones with the following probes,

Page 13, last full paragraph (lines 35-36)

~~Figure 11 shows~~ Figures 11A-C show the hybridisation of wheat DNA from chromosome-engineered lines using the following probes:

Page 18, third full paragraph (lines 14-27)

~~Figure 27 shows~~ Figures 27A-R show the results of transient expression assays typical of each promoter and target tissue. The photographs (40 x magnification) of representative tissue resulting from the transient expression assays typical of each promoter and target tissue revealed under a Leica microscope with blue light illumination. Photographs were taken 48 to 72 hours after tissue bombardment. The promoter constructs are listed as follows, (with the panels showing endosperm, embryo and leaf expression listed in respective order):
pact_jsgfp_nos (panels a,g and m); pwsssIpro1gfpNOT (panels b, h and n);
pwsssIpro2gfpNOT (panels c, i and o); psbeIIpro1gfpNOT (panels d, j and p);
psbeIIpro2gfpNOT (panels e, k and q); pZLgfpNOT (Panels f, l and r).